

Serial Number: 09434, 965**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☒ other invalid text, such as extraneous material at end of file.
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

1644

RAW SEQUENCE LISTING DATE: 11/02/2000
 PATENT APPLICATION: US/09/434,965 TIME: 12:37:24

Input Set : A:\Cpg.pto
 Output Set: N:\CRF3\11022000\I434965.raw

SEQUENCE LISTING

4 (1) GENERAL INFORMATION:

6 (i) APPLICANT: Luxembourg, Alain T.
 7 Jackson, Michael R.
 8 Peterson, Per A.
 10 (ii) TITLE OF INVENTION: PURIFICATION OF ANTIGEN-SPECIFIC T CELLS
 12 (iii) NUMBER OF SEQUENCES: 11
 14 (iv) CORRESPONDENCE ADDRESS:
 15 (A) ADDRESSEE: Johnson & Johnson
 16 (B) STREET: One Johnson & Johnson Plaza
 17 (C) CITY: New Brunswick
 18 (D) STATE: NJ
 19 (E) COUNTRY: USA
 20 (F) ZIP: 08933-7003
 22 (v) COMPUTER READABLE FORM:
 23 (A) MEDIUM TYPE: Floppy disk
 24 (B) COMPUTER: IBM PC compatible
 25 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 26 (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
 28 (vi) CURRENT APPLICATION DATA:

C--> 29 (A) APPLICATION NUMBER: US/09/434,965

C--> 30 (B) FILING DATE: 05-Nov-1999

31 (C) CLASSIFICATION:

33 (viii) ATTORNEY/AGENT INFORMATION:

34 (A) NAME: Wallen, John W.
 35 (B) REGISTRATION NUMBER: 35,403
 36 (C) REFERENCE/DOCKET NUMBER: ORT-841

38 (ix) TELECOMMUNICATION INFORMATION:

39 (A) TELEPHONE: 732-524-2806
 40 (B) TELEFAX: 732-524-2808

45 (2) INFORMATION FOR SEQ ID NO: 1:

47 (i) SEQUENCE CHARACTERISTICS:

48 (A) LENGTH: 9 amino acids
 49 (B) TYPE: amino acid
 50 (C) STRANDEDNESS: single
 51 (D) TOPOLOGY: linear

53 (ii) MOLECULE TYPE: peptide

55 (iv) ANTI-SENSE: NO

57 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

59 Gln Leu Ser Pro Phe Pro Phe Asp Leu
 60 1 5

63 (2) INFORMATION FOR SEQ ID NO: 2:

65 (i) SEQUENCE CHARACTERISTICS:

66 (A) LENGTH: 8 amino acids
 67 (B) TYPE: amino acid
 68 (C) STRANDEDNESS: single
 69 (D) TOPOLOGY: linear

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71      (ii) MOLECULE TYPE: peptide
73      (iv) ANTI-SENSE: NO
75      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
77      Leu Ser Pro Phe Pro Phe Asp Leu
78      1           5

81 (2) INFORMATION FOR SEQ ID NO: 3:
83      (i) SEQUENCE CHARACTERISTICS:
84          (A) LENGTH: 9 amino acids
85          (B) TYPE: amino acid
86          (C) STRANDEDNESS: single
87          (D) TOPOLOGY: linear
89      (ii) MOLECULE TYPE: peptide
91      (iv) ANTI-SENSE: NO
93      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
96      Ser Pro Phe Pro Phe Asp Leu Leu Leu
97      1           5

100 (2) INFORMATION FOR SEQ ID NO: 4:
102      (i) SEQUENCE CHARACTERISTICS:
103          (A) LENGTH: 8 amino acids
104          (B) TYPE: amino acid
105          (C) STRANDEDNESS: single
106          (D) TOPOLOGY: linear
108      (ii) MOLECULE TYPE: peptide
110      (iv) ANTI-SENSE: NO
112      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
114      Leu Ser Ala Phe Pro Phe Asp Leu
115      1           5

118 (2) INFORMATION FOR SEQ ID NO: 5:
120      (i) SEQUENCE CHARACTERISTICS:
121          (A) LENGTH: 8 amino acids
122          (B) TYPE: amino acid
123          (C) STRANDEDNESS: single
124          (D) TOPOLOGY: linear
126      (ii) MOLECULE TYPE: peptide
128      (iv) ANTI-SENSE: NO
130      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
132      Glu Gln Tyr Lys Phe Tyr Ser Val
133      1           5

136 (2) INFORMATION FOR SEQ ID NO: 6:
138      (i) SEQUENCE CHARACTERISTICS:
139          (A) LENGTH: 8 amino acids
140          (B) TYPE: amino acid
141          (C) STRANDEDNESS: single
142          (D) TOPOLOGY: linear
144      (ii) MOLECULE TYPE: peptide
147      (iv) ANTI-SENSE: NO
149      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
151      Ser Ile Tyr Arg Tyr Tyr Gly Leu
152      1           5

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RAW SEQUENCE LISTING
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155 (2) INFORMATION FOR SEQ ID NO: 7:
157   (i) SEQUENCE CHARACTERISTICS:
158       (A) LENGTH: 9 amino acids
159       (B) TYPE: amino acid
160       (C) STRANDEDNESS: single
161       (D) TOPOLOGY: linear
163   (ii) MOLECULE TYPE: peptide
165   (iv) ANTI-SENSE: NO
167   (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
169       Arg Pro Gln Ala Ser Gly Val Tyr Met
170       1           5
173 (2) INFORMATION FOR SEQ ID NO: 8:
175   (i) SEQUENCE CHARACTERISTICS:
176       (A) LENGTH: 9 amino acids
177       (B) TYPE: amino acid
178       (C) STRANDEDNESS: single
179       (D) TOPOLOGY: linear
181   (ii) MOLECULE TYPE: peptide
183   (iv) ANTI-SENSE: NO
185   (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
187       Tyr Pro His Phe Met Pro Thr Asn Leu
188       1           5
190 (2) INFORMATION FOR SEQ ID NO: 9:
192   (i) SEQUENCE CHARACTERISTICS:
193       (A) LENGTH: 8 amino acids
194       (B) TYPE: amino acid
195       (C) STRANDEDNESS: single
196       (D) TOPOLOGY: linear
198   (ii) MOLECULE TYPE: peptide
200   (iv) ANTI-SENSE: NO
202   (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
204       Ser Ile Ile Asn Phe Glu Lys Leu
205       1           5
208 (2) INFORMATION FOR SEQ ID NO: 10:
210   (i) SEQUENCE CHARACTERISTICS:
211       (A) LENGTH: 8 amino acids
212       (B) TYPE: amino acid
213       (C) STRANDEDNESS: single
214       (D) TOPOLOGY: linear
216   (ii) MOLECULE TYPE: peptide
218   (iv) ANTI-SENSE: NO
220   (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
222       Arg Gly Tyr Val Tyr Gln Gly Asn
223       1           5
226 (2) INFORMATION FOR SEQ ID NO: 11:
228   (i) SEQUENCE CHARACTERISTICS:
229       (A) LENGTH: 8 amino acids
230       (B) TYPE: amino acid
231       (C) STRANDEDNESS: single

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232 (D) TOPOLOGY: linear
234 (ii) MOLECULE TYPE: peptide
236 (iv) ANTI-SENSE: NO
238 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
240 Glu Ile Ile Asn Phe Glu Lys Leu
241 1 5

VERIFICATION SUMMARY DATE: 11/02/2000
PATENT APPLICATION: US/09/434,965 TIME: 12:37:25

Input Set : A:\Cpg.pto
Output Set: N:\CRF3\11022000\I434965.raw

L:29 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:30 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]